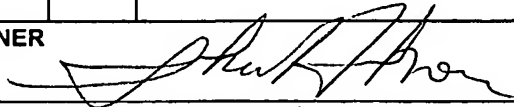
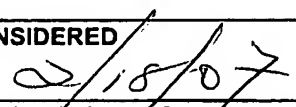


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INFORMATION DISCLOSURE CITATION (Use several sheets if necessary)				APPLICANT Hassan M. FATHALLAH-SHAYKH			
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<b>U.S. PATENT DOCUMENTS</b>							
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<b>FOREIGN PATENT DOCUMENTS</b>							
	REF	DOCUMENT NUMBER	DATE	COUNTRY	CLASS	SUB- CLASS	TRANSLATION YES NO
<b>OTHER DOCUMENTS (Including Author, Title, Date, Pertinent Pages, Etc.)</b>							
A		Fathallah-Shayk, H. M. et al., "Mathematical Modeling of Noise and Discovery of Genetic Expression Classes in Gliomas," <i>Oncogene</i> , October 17, 2002, Vol. 21, No. 47, pp. 7164-7174.					
A		Fathallah-Shaykh, H. M. et al., "Supplementary Information to Mathematical Modeling of Noise and Discovery of Genetic Expression Classes in Gliomas," published on the Web at <a href="http://www.rush.edu/neurosci/Fathallah.html">www.rush.edu/neurosci/Fathallah.html</a> , after October 17, 2002.					
A		Fathallah-Shaykh, H. M. et al., "Genomic Expression Discovery Predicts Pathways and Opposing Functions behind Phenotypes," <i>The Journal of Biological Chemistry</i> , Vol. 278, No. 26, June 27, 2003, pp. 23830-23833; published by The American Society for Biochemistry and Molecular Biology, Inc.					
A		Fathallah-Shaykh, H. M. et al., "Supplementary Information to Genomic Expression Discovery Predicts Pathways And Opposing Functions Behind Phenotypes," published on the Web at <a href="http://www.nature.com/onc/index.html">www.nature.com/onc/index.html</a> , April 16, 2003.					
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## FOREIGN PATENT DOCUMENTS

	REF	DOCUMENT NUMBER	DATE	COUNTRY	CLASS	SUB- CLASS	TRANSLATION	
							YES	NO

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<i>A</i>		Tusher, et al., "Significance Analysis of microarrays applied to the ionizing radiation response," <i>PNAS</i> , Vol. 98, No. 9, pp. 5116-5121, April 24, 2001.
<i>A</i>		Newton, et al., "On differential variability of expression ratios: improving statistical inference about gene expression changes from microarray data," <i>Journal of Computational Biology</i> , Vol. 8, No. 1, pp. 37-52, 2001.
<i>A</i>		Baggerly, et al., "Identifying differentially expressed genes in cDNA microarray experiments," <i>Journal of Computational Biology</i> , Vol. 8, No. 6, pp. 639-659, 2001.
<i>A</i>		Alter, et al., "Singular value decomposition for genome-wide expression data processing and modeling," <i>PNAS</i> , Vol. 97, No. 18, pp. 10101-10106, August 29, 2000.

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